IN THE SPECIFICATION:

After page 30, please insert the attached Sequence Listing.

Please replace paragraph on page 16, starting at line 1 and ending at line 2 with the following replacement paragraph:

end downstream the EcoRI site gaattc (SEQ ID NO:1), i.e. the lack of the 3' end from nucleotide 2014 included.

Please replace paragraph on page 16, starting at line 18 and ending at line 26 with the following replacement paragraph:

To fuse the *STM* coding region to the engrailed repressor domain the frame of the unique XbaI cloning site in pRT Ωeng had to be shifted which was achieved by inserting a GATCTCGA (SEQ ID NO:2) adapter into the upstream BamHI site, which was destroyed. The STM coding region previously amplified by reverse transcriptase PCR from Arabidopsis RNA with an 5' terminal XbaI (upstream the translation start at ATG) was inserted into the XbaI site of the adpted pRT Ωeng. The STM coding sequences used contain a BamHI site immediately preceding the natural translation stop codon, which was used subsequently to create the c-terminal GR fusion.

REMARKS

In response to the Notice to Comply dated October 4, 2001, Applicant submits herewith a Sequence Listing in paper and computer readable form for the above-identified application and requests that the sequence listing be inserted into the specification as indicated above. Applicant also submits herewith a copy of the Notice to Comply.

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